Ian R Gizer

List of Publications by Year in descending order

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	394421	223800
2,236	19	46
citations	h-index	g-index
57	57	3442
docs citations	times ranked	citing authors
	citations 57	2,236 19 citations h-index 57 57

#	Article	IF	CITATIONS
1	Candidate gene studies of ADHD: a meta-analytic review. Human Genetics, 2009, 126, 51-90.	3.8	871
2	The genetics of attention deficit hyperactivity disorder. Clinical Psychology Review, 2006, 26, 396-432.	11.4	136
3	A polymorphism in the norepinephrine transporter gene alters promoter activity and is associated with attention-deficit hyperactivity disorder. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 19164-19169.	7.1	131
4	Cannabis dependence in the San Francisco Family Study: Age of onset of use, DSM-IV symptoms, withdrawal, and heritability. Addictive Behaviors, 2010, 35, 102-110.	3.0	89
5	Measuring historical trauma in an American Indian community sample: Contributions of substance dependence, affective disorder, conduct disorder and PTSD. Drug and Alcohol Dependence, 2013, 133, 180-187.	3.2	82
6	Evidence for a Genetic Component for Substance Dependence in Native Americans. American Journal of Psychiatry, 2013, 170, 154-164.	7.2	62
7	Choline transporter gene variation is associated with attention-deficit hyperactivity disorder. Journal of Neurodevelopmental Disorders, 2009, 1, 252-263.	3.1	61
8	Lifetime history of traumatic events in an American Indian community sample: Heritability and relation to substance dependence, affective disorder, conduct disorder and PTSD. Journal of Psychiatric Research, 2013, 47, 155-161.	3.1	58
9	ADH and ALDH Polymorphisms and Alcohol Dependence in Mexican and Native Americans. American Journal of Drug and Alcohol Abuse, 2012, 38, 389-394.	2.1	51
10	The adrenergic receptor Â-A gene (ADRA2A) and neuropsychological executive functions as putative endophenotypes for childhood ADHD. Cognitive, Affective and Behavioral Neuroscience, 2006, 6, 18-30.	2.0	49
11	Association of Alcohol Dehydrogenase Genes with Alcohol-Related Phenotypes in a Native American Community Sample. Alcoholism: Clinical and Experimental Research, 2011, 35, 2008-2018.	2.4	34
12	Variant calling in low-coverage whole genome sequencing of a Native American population sample. BMC Genomics, 2014, 15, 85.	2.8	33
13	GENETIC STUDY: Heritability and a genomeâ€wide linkage analysis of a Type II/B cluster construct for cannabis dependence in an American Indian community. Addiction Biology, 2009, 14, 338-348.	2.6	32
14	Age at Regular Drinking, Clinical Course, and Heritability of Alcohol Dependence in the San Francisco Family Study: A Gender Analysis. American Journal on Addictions, 2010, 19, 101-110.	1.4	32
15	Relations between multi-informant assessments of ADHD symptoms, DAT1, and DRD4 Journal of Abnormal Psychology, 2008, 117, 869-880.	1.9	29
16	Double dissociation between lab measures of inattention and impulsivity and the dopamine transporter gene (DAT1) and dopamine D4 receptor gene (DRD4) Journal of Abnormal Psychology, 2012, 121, 1011-1023.	1.9	29
17	EEG spectral phenotypes: Heritability and association with marijuana and alcohol dependence in an American Indian community study. Drug and Alcohol Dependence, 2010, 106, 101-110.	3.2	26
18	Linkage analyses of cannabis dependence, craving, and withdrawal in the San Francisco family study. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 802-811.	1.7	21

#	Article	IF	Citations
19	Item Response Theory Analysis of Binge Drinking and Its Relationship to Lifetime Alcohol Use Disorder Symptom Severity in an American Indian Community Sample. Alcoholism: Clinical and Experimental Research, 2011, 35, 984-995.	2.4	21
20	Metaâ€Analysis of Genetic Influences on Initial Alcohol Sensitivity. Alcoholism: Clinical and Experimental Research, 2018, 42, 2349-2359.	2.4	21
21	Indexing the â€~dark side of addiction': substanceâ€induced affective symptoms and alcohol use disorders. Addiction, 2019, 114, 139-149.	3.3	21
22	Genome-wide scan for self-rating of the effects of alcohol in American Indians. Psychiatric Genetics, 2010, 20, 221-228.	1.1	20
23	Linkage analyses of stimulant dependence, craving, and heavy use in American Indians. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 772-780.	1.7	20
24	EEG alpha phenotypes: linkage analyses and relation to alcohol dependence in an American Indian community study. BMC Medical Genetics, 2010, 11, 43.	2.1	19
25	Genetic loci for alcohol-related life events and substance-induced affective symptoms: indexing the $\hat{a} \in \mathbb{R}$ and substance-induced affective symptoms: indexing the $\hat{a} \in \mathbb{R}$ and $\hat{a} \in \mathbb{R}$ of addiction. Translational Psychiatry, 2019, 9, 71.	4.8	19
26	Common genetic contributions to high-risk trauma exposure and self-injurious thoughts and behaviors. Psychological Medicine, 2019, 49, 421-430.	4.5	18
27	Association and ancestry analysis of sequence variants in ADH and ALDH using alcoholâ€related phenotypes in a Native American community sample. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 673-683.	1.7	17
28	Comorbidity of Alcohol Use Disorder and Chronic Pain: Genetic Influences on Brain Reward and Stress Systems. Alcoholism: Clinical and Experimental Research, 2017, 41, 1831-1848.	2.4	17
29	Whole genome sequence study of cannabis dependence in two independent cohorts. Addiction Biology, 2018, 23, 461-473.	2.6	16
30	Linkage scan of alcohol dependence in the UCSF Family Alcoholism Study. Drug and Alcohol Dependence, 2011, 113, 125-132.	3.2	15
31	Item Response Theory analyses of DSM-IV and DSM-5 stimulant use disorder criteria in an American Indian community sample. Drug and Alcohol Dependence, 2014, 135, 29-36.	3.2	15
32	Genomeâ€wide metaâ€analysis identifies a novel susceptibility signal at <i>CACNA2D3</i> for nicotine dependence. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 557-567.	1.7	15
33	<i>CHRNA5</i> and <i>CHRNA3</i> Variants and Level of Neuroticism in Young Adult Mexican American Men and Women. Twin Research and Human Genetics, 2014, 17, 80-88.	0.6	13
34	Correlation analysis of genetic admixture and social identification with body mass index in a Native American Community. American Journal of Human Biology, 2014, 26, 347-360.	1.6	13
35	A cisâ€ <scp>eQTL</scp> in <i><scp>OPRM</scp>1</i> is Associated with Subjective Response to Alcohol and Alcohol Use. Alcoholism: Clinical and Experimental Research, 2017, 41, 929-938.	2.4	13
36	Protective variant associated with alcohol dependence in a Mexican American cohort. BMC Medical Genetics, 2014, 15, 136.	2.1	12

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37	Single nucleotide polymorphisms in the <i><scp>REG TNNA2</scp></i> region of chromosome 2 and <i><scp>NEIL3</scp></i> associated with impulsivity in a <scp>N</scp> ative <scp>A</scp> merican sample. Genes, Brain and Behavior, 2016, 15, 568-577.	2.2	12
38	Genetic variation in the exome: Associations with alcohol and tobacco co-use Psychology of Addictive Behaviors, 2017, 31, 354-366.	2.1	10
39	Heritability of MMPI-2 Scales in the UCSF Family Alcoholism Study. Journal of Addictive Diseases, 2010, 29, 84-97.	1.3	7
40	Genome-wide linkage scan of antisocial behavior, depression, and impulsive substance use in the UCSF family alcoholism study. Psychiatric Genetics, 2012, 22, 235-244.	1.1	7
41	Event-related oscillations to affective stimuli: Heritability, linkage and relationship to externalizing disorders. Journal of Psychiatric Research, 2012, 46, 256-263.	3.1	7
42	Contributions of Ethnicity to Differential Item Functioning of Cannabis Abuse and Dependence Symptoms. Journal of Studies on Alcohol and Drugs, 2013, 74, 320-328.	1.0	7
43	Stimulant Dependence and Stimulant-Associated Psychosis. Journal of Addiction Medicine, 2014, 8, 241-248.	2.6	7
44	Polygenic risk scores for cigarettes smoked per day do not generalize to a Native American population. Drug and Alcohol Dependence, 2016, 167, 95-102.	3.2	7
45	Associations Between Genomic Variants in Alcohol Dehydrogenase Genes and Alcohol Symptomatology in American Indians and European Americans: Distinctions and Convergence. Alcoholism: Clinical and Experimental Research, 2017, 41, 1695-1704.	2.4	7
46	Genetic Influences on Evening Preference Overlap with Those for Bipolar Disorder in a Sample of Mexican Americans and American Indians. Twin Research and Human Genetics, 2017, 20, 499-510.	0.6	7
47	Molecular genetic approaches to understanding the comorbidity of psychiatric disorders. Development and Psychopathology, 2016, 28, 1089-1101.	2.3	6
48	Linkage scan of nicotine dependence in the University of California, San Francisco (UCSF) Family Alcoholism Study. Psychological Medicine, 2011, 41, 799-808.	4.5	4
49	Genome-Wide Association Studies of Substance Use: Considerations Regarding Populations and Phenotypes. Biological Psychiatry, 2015, 77, 423-424.	1.3	4
50	A Novel Tobacco Use Phenotype Suggests the 15q25 and 19q13 Loci May be Differentially Associated With Cigarettes per Day and Tobacco-Related Problems. Nicotine and Tobacco Research, 2016, 19, ntw260.	2.6	4
51	Effects of genetic risk for alcohol dependence and onset of regular drinking on the progression to alcohol dependence: A polygenic risk score approach. Drug and Alcohol Dependence, 2022, 230, 109117.	3.2	4
52	Variants Near CCK Receptors are Associated With Electrophysiological Responses to Pre-pulse Startle Stimuli in a Mexican American Cohort. Twin Research and Human Genetics, 2015, 18, 727-737.	0.6	2
53	Differential item functioning analysis of the CUDIT and relations with alcohol and tobacco use among men across five ethnic groups: The HELIUS study Psychology of Addictive Behaviors, 2019, 33, 697-709.	2.1	2
54	DISC1 loci not associated with anhedonia in individuals with genetic liability for schizophrenia. Psychiatric Genetics, 2014, 24, 120-121.	1.1	1

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55	Molecular Genetic Approaches to Studying the Externalizing Spectrum. , 2015, , .		0
56	Molecular Genetics of the Externalizing Spectrum. , 2015, , .		0
57	Association of Predicted Expression and Multimodel Association Analysis of Substance Abuse Traits. Complex Psychiatry, 2022, 8, 35-46.	0.9	0